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RAW SEQUENCE LISTING PATENT APPLICATION US/08/882,435

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This Raw Listing contains the General Information Section and up to the first 5 pages.

SEQUENCE LISTING 1 ENTERED 2 (1) General Information: 3 (i) APPLICANT: Hoxie, James A. 5 6 (ii) TITLE OF INVENTION: ANTIBODIES DIRECTED AGAINST CELLULAR 8 CORECEPTORS FOR HUMAN IMMUNODEFICIENCY VIRUS AND METHODS OF USING THE SAME 9 10 11 (iii) NUMBER OF SEQUENCES: 6 12 13 (iv) CORRESPONDENCE ADDRESS: (A) ADDRESSEE: PANITCH SCHWARZE JACOBS & NADEL, P.C. 14 (B) STREET: One Commerce Square, 2005 Market Street, 22nd Floor 15 16 (C) CITY: Philadelphia 17 (D) STATE: Pennsylvania 18 (E) COUNTRY: U.S. (F) ZIP: 19103-7086 19 20 (v) COMPUTER READABLE FORM: 21 (A) MEDIUM TYPE: Floppy disk 22 (B) COMPUTER: IBM PC compatible 23 (C) OPERATING SYSTEM: PC-DOS/MS-DOS 24 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 25 26 (vi) CURRENT APPLICATION DATA: 27 (A) APPLICATION NUMBER: US 08/882,435 28 29 (B) FILING DATE: 25-JUN-1997 (C) CLASSIFICATION: 30 31 32 (viii) ATTORNEY/AGENT INFORMATION: 33 (A) NAME: Doyle Leary, Ph.D., Kathryn (B) REGISTRATION NUMBER: 36,317 34 (C) REFERENCE/DOCKET NUMBER: 9596-11U1 35 36 (ix) TELECOMMUNICATION INFORMATION: 37 (A) TELEPHONE: 215-965-1284 38 (B) TELEFAX: 215-567-2991 39 40 (C) TELEX: 831-494 41 42 (2) INFORMATION FOR SEQ ID NO:1: 43 44 45 (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 27 base pairs

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			IIII OI DEII. DEFSONIUN
47 48 49 50		(B) TYPE: nucleic acid(C) STRANDEDNESS: double(D) TOPOLOGY: linear	
51	(ii)	MOLECULE TYPE: DNA (genomic)	
52 53	(iii)	HYPOTHETICAL: NO	
54 55 56 57 58	(iv)	ANTI-SENSE: NO	
59 60	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:1:	
61 62	TAGATATC	TT ACCATGGAGG GGATCAG	27
63 64	(2) INFO	RMATION FOR SEQ ID NO:2:	
65 66 67 68 69 70	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear	
71 72 73	(ii)	MOLECULE TYPE: DNA (genomic)	
74	(iii)	HYPOTHETICAL: NO	
75 76 77 78 79	(iv)	ANTI-SENSE: NO	
80 81 82	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:	
83	TAGCGGCG	CT TAGTGGAGTG AAAACTTG	28
84 85	(2) INFO	RMATION FOR SEQ ID NO:3:	
86 87 88 89 90 91	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear	
93 94	(ii)	MOLECULE TYPE: DNA (genomic)	
95 96	(iii)	HYPOTHETICAL: NO	
97 98 99	(iv)	ANTI-SENSE: NO	

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	INFO) 1 SE1: S2930.raw			
100					
101					
102	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:				
103					
104	TGGATATTGC CATCAATGAC C 21				
105					
106	(2) INFORMATION FOR SEQ ID NO:4:				
107					
108	(i) SEQUENCE CHARACTERISTICS:				
109	(A) LENGTH: 22 base pairs				
110	(B) TYPE: nucleic acid				
111	(C) STRANDEDNESS: double				
112	(D) TOPOLOGY: linear				
113					
114	(ii) MOLECULE TYPE: DNA (genomic)				
115					
116	(iii) HYPOTHETICAL: NO				
117					
118	(iv) ANTI-SENSE: NO				
119					
120					
121					
122					
123	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:				
124					
125	GATGGCATGG ACTGTGGTCA TG	22			
126					
127	(2) INFORMATION FOR SEQ ID NO:5:				
128					
129	(i) SEQUENCE CHARACTERISTICS:				
130	(A) LENGTH: 48 base pairs				
131	(B) TYPE: nucleic acid				
132	(C) STRANDEDNESS: double				
133	(D) TOPOLOGY: linear				
134					
135	(ii) MOLECULE TYPE: DNA (genomic)				
136					
137	(iii) HYPOTHETICAL: NO				
138					
139	(iv) ANTI-SENSE: NO				
140					
141					
142					
143					
144	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:				
145					
146	GGCTCATCCG GTCGACGAAT CAGACAAGTG AGTATGAAGG GTAGTAAG 48				
147					
148	(2) INFORMATION FOR SEQ ID NO:6:				
149					
150	(i) SEQUENCE CHARACTERISTICS:				
151	(A) LENGTH: 34 base pairs				
152	(B) TYPE: nucleic acid				

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		1141 O1 3E1. 32730.1W
153	(C) STRANDEDNESS: double	
154	(D) TOPOLOGY: linear	
155		
156	(ii) MOLECULE TYPE: DNA (genomic)	
157		
158	(iii) HYPOTHETICAL: NO	
159		
160	(iv) ANTI-SENSE: NO	
161		
162		
163		
164		
165	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
166		
167	CTGCTGATAT CGCTGTCCCT CACAGGAGGG CGAG	34
168		

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